



## **RAW SEQUENCE LISTING** **ERROR REPORT**

RECEIVED

AUG 27 2002

TECH CENTER 1600/2900

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/613,006  
Source: 1600  
Date Processed by STIC: 8/22/02

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**

**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

**FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.**

**FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.**

**PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)**

**PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)**

**TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:**

**<http://www.uspto.gov/web/offices/pac/checker>**

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:  
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202  
Or  
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

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Does Not Comply  
Corrected Diskette Needed

1600

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/613,006

DATE: 08/22/2002

TIME: 15:12:18

*Errors on PR3+5*

Input Set : A:\52949-20001.00.txt

Output Set: N:\CRF4\08222002\I613006.raw

```

4 <110> APPLICANT: Schena, Mark A.
6 <120> TITLE OF INVENTION: MICROARRAY METHOD OF GENOTYPING MULTIPLE
7   SAMPLES AT MULTIPLE LOCI
9 <130> FILE REFERENCE: 529492000100
11 <140> CURRENT APPLICATION NUMBER: US 09/613,006
12 <141> CURRENT FILING DATE: 2000-07-10
14 <160> NUMBER OF SEQ ID NOS: 30
16 <170> SOFTWARE: FastSEQ for Windows Version 4.0
18 <210> SEQ ID NO: 1
19 <211> LENGTH: 20
20 <212> TYPE: DNA
21 <213> ORGANISM: Homo sapiens
23 <220> FEATURE:
24 <221> NAME/KEY: misc_feature
25 <222> LOCATION: 1
26 <223> OTHER INFORMATION: n = adenine with C6 modification
28 <400> SEQUENCE: 1
W--> 29 naacagacac catggtgcac                                20
31 <210> SEQ ID NO: 2
32 <211> LENGTH: 20
33 <212> TYPE: DNA
34 <213> ORGANISM: Homo sapiens
36 <220> FEATURE:
37 <221> NAME/KEY: misc_feature
38 <222> LOCATION: 1
39 <223> OTHER INFORMATION: n = cytosine with a C6 modification
41 <400> SEQUENCE: 2
W--> 42 nccacagggc agtaacggca                                20
44 <210> SEQ ID NO: 3
45 <211> LENGTH: 20
46 <212> TYPE: DNA
47 <213> ORGANISM: Homo sapiens
49 <220> FEATURE:
50 <221> NAME/KEY: misc_feature
51 <222> LOCATION: 1
52 <223> OTHER INFORMATION: n = guanine with a C6 modification
54 <400> SEQUENCE: 3
W--> 55 ncaaggtgaa cgtggatgaa                                20
57 <210> SEQ ID NO: 4
58 <211> LENGTH: 20
59 <212> TYPE: DNA
60 <213> ORGANISM: Homo sapiens
62 <220> FEATURE:

```

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```

63 <221> NAME/KEY: misc_feature
64 <222> LOCATION: 1
65 <223> OTHER INFORMATION: n = guanine with a C6 modification
67 <400> SEQUENCE: 4
W--> 68 ntaaccttga taccaacctg                20
70 <210> SEQ ID NO: 5
71 <211> LENGTH: 20
72 <212> TYPE: DNA
73 <213> ORGANISM: Homo sapiens
75 <220> FEATURE:
76 <221> NAME/KEY: misc_feature
77 <222> LOCATION: 1
78 <223> OTHER INFORMATION: n = cytosine with a C6 modification
80 <400> SEQUENCE: 5
W--> 81 ntggcaccat taaagaaaat                20
83 <210> SEQ ID NO: 6
84 <211> LENGTH: 20
85 <212> TYPE: DNA
86 <213> ORGANISM: Homo sapiens
88 <220> FEATURE:
89 <221> NAME/KEY: misc_feature
90 <222> LOCATION: 1
91 <223> OTHER INFORMATION: n = thymine with a C6 modification
93 <400> SEQUENCE: 6
W--> 94 ntctgtatct atattcatca                20
96 <210> SEQ ID NO: 7
97 <211> LENGTH: 20
98 <212> TYPE: DNA
99 <213> ORGANISM: Homo sapiens
101 <220> FEATURE:
102 <221> NAME/KEY: misc_feature
103 <222> LOCATION: 1
104 <223> OTHER INFORMATION: n = thymine with a C6 modification
106 <400> SEQUENCE: 7
W--> 107 ngggctgttc taacccccac                20
109 <210> SEQ ID NO: 8
110 <211> LENGTH: 20
111 <212> TYPE: DNA
112 <213> ORGANISM: Homo sapiens
114 <220> FEATURE:
115 <221> NAME/KEY: misc_feature
116 <222> LOCATION: 1
117 <223> OTHER INFORMATION: n = adenine with a C6 modification
119 <400> SEQUENCE: 8
W--> 120 naccactgg agcccctgac                20
122 <210> SEQ ID NO: 9
123 <211> LENGTH: 20
124 <212> TYPE: DNA
125 <213> ORGANISM: Homo sapiens

```

## RAW SEQUENCE LISTING

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Output Set: N:\CRF4\08222002\I613006.raw

```

127 <220> FEATURE:
128 <221> NAME/KEY: misc_feature
129 <222> LOCATION: 1
130 <223> OTHER INFORMATION: n = cytosine with a C6 modification
132 <400> SEQUENCE: 9
W--> 133 ncacaggatc agaggctggg 20
135 <210> SEQ ID NO: 10
136 <211> LENGTH: 20
137 <212> TYPE: DNA
138 <213> ORGANISM: Homo sapiens
140 <220> FEATURE:
141 <221> NAME/KEY: misc_feature
142 <222> LOCATION: 1
143 <223> OTHER INFORMATION: n = guanine with a C6 modification
145 <400> SEQUENCE: 10
W--> 146 ngtagtaatg agcgtgcagc 20
148 <210> SEQ ID NO: 11
149 <211> LENGTH: 15
150 <212> TYPE: DNA
151 <213> ORGANISM: Homo sapiens
153 <220> FEATURE:
154 <221> NAME/KEY: misc_feature
155 <222> LOCATION: 1
156 <223> OTHER INFORMATION: n = guanine with a Cy3 label
158 <221> NAME/KEY: variation
159 <222> LOCATION: 9
160 <223> OTHER INFORMATION: n = a or t
W--> 162 <400> 11
W--> 163 nactcctgng gagaa 15
165 <210> SEQ ID NO: 12
166 <211> LENGTH: 15
167 <212> TYPE: DNA
168 <213> ORGANISM: Homo sapiens
170 <220> FEATURE:
171 <221> NAME/KEY: misc_feature
172 <222> LOCATION: 1
173 <223> OTHER INFORMATION: n = guanine with Cy5 labels
175 <400> SEQUENCE: 12
W--> 176 nactcctang gagaa 15
178 <210> SEQ ID NO: 13
179 <211> LENGTH: 15
180 <212> TYPE: DNA
181 <213> ORGANISM: Homo sapiens
183 <220> FEATURE:
184 <221> NAME/KEY: misc_feature
185 <222> LOCATION: 1
186 <223> OTHER INFORMATION: n = thymine attached to a Cy5 label
188 <400> SEQUENCE: 13
W--> 189 nggtggtgag gccct 15

```

*→ n also found at location 9*

## RAW SEQUENCE LISTING

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DATE: 08/22/2002

TIME: 15:12:18

Input Set : A:\52949-20001.00.txt

Output Set: N:\CRF4\08222002\I613006.raw

```

191 <210> SEQ ID NO: 14
192 <211> LENGTH: 15
193 <212> TYPE: DNA
194 <213> ORGANISM: Homo sapiens
196 <220> FEATURE:
197 <221> NAME/KEY: misc_feature
198 <222> LOCATION: 1
199 <223> OTHER INFORMATION: n = thymine attached to a Cy3 label
201 <400> SEQUENCE: 14
W--> 202 nggtggtgaag gccct 15
204 <210> SEQ ID NO: 15
205 <211> LENGTH: 15
206 <212> TYPE: DNA
207 <213> ORGANISM: Homo sapiens
209 <220> FEATURE:
210 <221> NAME/KEY: misc_feature
211 <222> LOCATION: 1
212 <223> OTHER INFORMATION: n = adenine attached to Cy3
214 <400> SEQUENCE: 15
W--> 215 ntcatcttttg gtgtt 15
217 <210> SEQ ID NO: 16
218 <211> LENGTH: 15
219 <212> TYPE: DNA
220 <213> ORGANISM: Homo sapiens
222 <220> FEATURE:
223 <221> NAME/KEY: misc_feature
224 <222> LOCATION: 1
225 <223> OTHER INFORMATION: n = thymine attached to a Cy5 label
227 <400> SEQUENCE: 16
W--> 228 natcatcggt gtttc 15
230 <210> SEQ ID NO: 17
231 <211> LENGTH: 15
232 <212> TYPE: DNA
233 <213> ORGANISM: Homo sapiens
235 <220> FEATURE:
236 <221> NAME/KEY: misc_feature
237 <222> LOCATION: 1
238 <223> OTHER INFORMATION: n = cytosine attached to a Cy5 label
240 <400> SEQUENCE: 17
W--> 241 nactgccagg taagg 15
243 <210> SEQ ID NO: 18
244 <211> LENGTH: 15
245 <212> TYPE: DNA
246 <213> ORGANISM: Homo sapiens
248 <220> FEATURE:
249 <221> NAME/KEY: misc_feature
250 <222> LOCATION: 1
251 <223> OTHER INFORMATION: n = cytosine attached to a Cy3 label
253 <400> SEQUENCE: 18

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## RAW SEQUENCE LISTING

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PATENT APPLICATION: US/09/613,006

TIME: 15:12:18

Input Set : A:\52949-20001.00.txt

Output Set: N:\CRF4\08222002\I613006.raw

```

W--> 254 nactgccggg taagg 15
256 <210> SEQ ID NO: 19
257 <211> LENGTH: 15
258 <212> TYPE: DNA
259 <213> ORGANISM: Homo sapiens
261 <220> FEATURE:
262 <221> NAME/KEY: misc_feature
263 <222> LOCATION: 1
264 <223> OTHER INFORMATION: n = cytosine attached to a Cy3 label
266 <400> SEQUENCE: 19

W--> 267 naactggaac cattg 15
269 <210> SEQ ID NO: 20
270 <211> LENGTH: 15
271 <212> TYPE: DNA
272 <213> ORGANISM: Homo sapiens
274 <220> FEATURE:
275 <221> NAME/KEY: misc_feature
276 <222> LOCATION: 1
277 <223> OTHER INFORMATION: n = cytosine attached to a Cy5 label
279 <400> SEQUENCE: 20

W--> 280 naactgggac cattg 15
282 <210> SEQ ID NO: 21
283 <211> LENGTH: 15
284 <212> TYPE: DNA
285 <213> ORGANISM: Homo sapiens
287 <220> FEATURE:
288 <221> NAME/KEY: misc_feature
289 <222> LOCATION: 1
290 <223> OTHER INFORMATION: n = guanine attached to biotin
292 <221> NAME/KEY: variation
293 <222> LOCATION: 10 → n found at 9, but not at 10
294 <223> OTHER INFORMATION: n = a or t

W--> 296 <400> 21

W--> 297 nactcctgng gagaa 15
299 <210> SEQ ID NO: 22
300 <211> LENGTH: 15
301 <212> TYPE: DNA
302 <213> ORGANISM: Homo sapiens
304 <220> FEATURE:
305 <221> NAME/KEY: misc_feature
306 <222> LOCATION: 1
307 <223> OTHER INFORMATION: n = thymine attached to biotin
309 <400> SEQUENCE: 22

W--> 310 nggtgtaag gccct 15
312 <210> SEQ ID NO: 23
313 <211> LENGTH: 15
314 <212> TYPE: DNA
315 <213> ORGANISM: Homo sapiens
317 <220> FEATURE:

```

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/613,006

DATE: 08/22/2002  
TIME: 15:12:19

Input Set : A:\52949-20001.00.txt  
Output Set: N:\CRF4\08222002\I613006.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 1  
Seq#:2; N Pos. 1  
Seq#:3; N Pos. 1  
Seq#:4; N Pos. 1  
Seq#:5; N Pos. 1  
Seq#:6; N Pos. 1  
Seq#:7; N Pos. 1  
Seq#:8; N Pos. 1  
Seq#:9; N Pos. 1  
Seq#:10; N Pos. 1  
Seq#:11; N Pos. 1,9  
Seq#:12; N Pos. 1,9  
Seq#:13; N Pos. 1  
Seq#:14; N Pos. 1  
Seq#:15; N Pos. 1  
Seq#:16; N Pos. 1  
Seq#:17; N Pos. 1  
Seq#:18; N Pos. 1  
Seq#:19; N Pos. 1  
Seq#:20; N Pos. 1  
Seq#:21; N Pos. 1,9  
Seq#:22; N Pos. 1  
Seq#:23; N Pos. 1  
Seq#:24; N Pos. 1  
Seq#:25; N Pos. 1  
Seq#:26; N Pos. 1,9  
Seq#:27; N Pos. 1  
Seq#:28; N Pos. 1  
Seq#:29; N Pos. 1  
Seq#:30; N Pos. 1

## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/613,006

DATE: 08/22/2002

TIME: 15:12:19

Input Set : A:\52949-20001.00.txt

Output Set: N:\CRF4\08222002\I613006.raw

L:29 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0  
L:42 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0  
L:55 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0  
L:68 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0  
L:81 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0  
L:94 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0  
L:107 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0  
L:120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0  
L:133 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0  
L:146 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0  
L:162 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:11  
L:163 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0  
L:176 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0  
L:189 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0  
L:202 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0  
L:215 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0  
L:228 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0  
L:241 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0  
L:254 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0  
L:267 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0  
L:280 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0  
L:296 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:21  
L:297 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0  
L:310 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0  
L:323 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0  
L:336 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0  
L:349 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0  
L:365 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:26  
L:366 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0  
L:379 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0  
L:392 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0  
L:405 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0  
L:418 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0